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HIV RNA, CD4+ Percentage, and Risk of Hepatocellular Carcinoma by Cirrhosis Status

Jessie Torgersen , Michael J. Kallan, Dena M. Carbonari , Lesley S. Park , Rajni L. Mehta, Kathryn D'Addeo, Janet P. Tate, Joseph K. Lim, Matthew Bidwell Goetz, Maria C. Rodriguez-Barradas, Cynthia L. Gibert, Norbert Bräu, Sheldon T. Brown, Jason A. Roy, Tamar H. Taddei, Amy C. Justice , Vincent Lo Re III

See the Notes section for the authors' affiliations.

Correspondence to: Vincent Lo Re III, MD, MSCE, Division of Infectious Diseases, Department of Medicine, Center for Clinical Epidemiology and Biostatistics, Perelman School of Medicine, University of Pennsylvania, 836 Blockley Hall, 423 Guardian Dr, Philadelphia, PA 19104-6021 (e-mail: vincentl@penmedicine.upenn.edu).

Abstract

Background: Despite increasing incidence of hepatocellular carcinoma (HCC) among HIV-infected patients, it remains unclear if HIV-related factors contribute to development of HCC. We examined if higher or prolonged HIV viremia and lower CD4+ cell percentage were associated with HCC.

Methods: We conducted a cohort study of HIV-infected individuals who had HIV RNA, CD4+, and CD8+ cell counts and percentages assessed in the Veterans Aging Cohort Study (1999–2015). HCC was ascertained using Veterans Health Administration cancer registries and electronic records. Cox regression was used to determine hazard ratios (HR, 95% confidence interval [CI]) of HCC associated with higher current HIV RNA, longer duration of detectable HIV viremia (≥ 500 copies/mL), and current CD4+ cell percentage less than 14%, adjusting for traditional HCC risk factors. Analyses were stratified by previously validated diagnoses of cirrhosis prior to start of follow-up.

Results: Among 35 659 HIV-infected patients, 302 (0.8%) developed HCC over 281 441 person-years (incidence rate = 107.3 per 100 000 person-years). Among patients without baseline cirrhosis, higher HIV RNA (HR = 1.25, 95% CI = 1.12 to 1.40, per 1.0 \log_{10} copies/mL) and 12 or more months of detectable HIV (HR = 1.47, 95% CI = 1.02 to 2.11) were independently associated with higher risk of HCC. CD4+ percentage less than 14% was not associated with HCC in any model. Hepatitis C coinfection was a statistically significant predictor of HCC regardless of baseline cirrhosis status.

Conclusion: Among HIV-infected patients without baseline cirrhosis, higher HIV RNA and longer duration of HIV viremia increased risk of HCC, independent of traditional HCC risk factors. This is the strongest evidence to date that HIV viremia contributes to risk of HCC in this group.

Hepatocellular carcinoma (HCC) is a growing cause of cancer death among people living with HIV infection (1). Driven largely by hepatitis C virus (HCV) coinfection, hepatitis B virus (HBV) coinfection, and alcoholic liver disease, the incidence of HCC among HIV-infected persons in North America has risen more than fourfold from 1995 to 2009 (2). Moreover, HIV-infected individuals have a fourfold higher risk of HCC than uninfected persons (3).

Despite the rising incidence of HCC among HIV-infected individuals, the determinants of this malignancy remain largely unknown in this group (4). Three prior studies found no association between HIV suppression and risk of HCC (5–7), but these studies did not evaluate the effects of longer durations or higher levels of HIV viremia, nor did they account for cirrhosis. Moreover, previous studies among predominantly HIV and HCV-coinfected patients reported that lower absolute CD4+ cell

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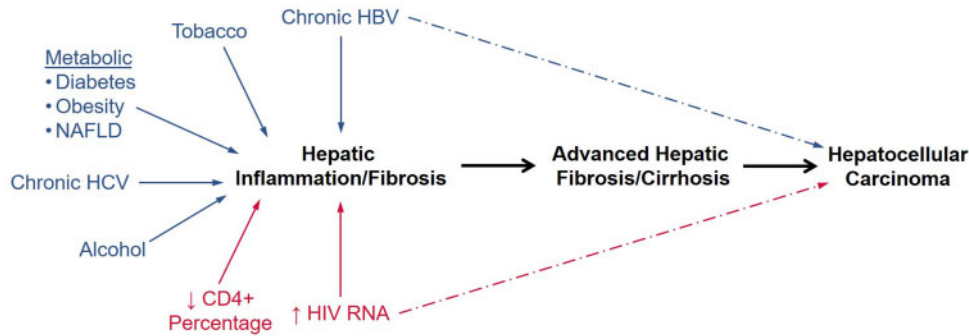


Figure 1. Pathway to development of hepatocellular carcinoma (HCC). The figure shows the contributions of the key modifiable determinants of HCC, including traditional risk factors (blue) and the hypothesized HIV-related determinants in this study (red). Both chronic hepatitis B virus (HBV) infection and HIV viremia could increase the risk of HCC by inducing hepatic fibrosis and cirrhosis or by directly promoting development of HCC outside of the liver fibrosis pathway. NAFLD = non-alcoholic fatty liver disease; RNA = ribonucleic acid.

counts increased the risk of HCC (6,8–12). However, absolute CD4+ cell count may decrease during cirrhosis because of portal hypertension-induced splenic sequestration (13). Thus, studies evaluating associations between absolute CD4+ cell count and HCC risk cannot determine whether observed associations are driven by HIV-related immunosuppression or progression of liver disease. Consequently, it remains unclear if higher HIV RNA levels, longer duration of detectable HIV, and HIV-related immunosuppression contribute to development of HCC independent of traditional determinants. Identifying such factors could help define the mechanisms for the high rate of HCC among HIV-infected persons.

Cirrhosis, which represents the late stage of progressive hepatic fibrosis due to chronic liver disease, is an important step in the causal pathway toward HCC (Figure 1) (14). Cirrhosis promotes HCC through telomere dysfunction and alterations of the liver milieu (eg, increased production of toxic hepatic metabolites, cytokines, growth factors, and products of oxidative stress) (15). Because cirrhosis increases the risk of HCC substantially, studies evaluating the factors associated with HCC must account for baseline cirrhosis status.

We evaluated HIV-related and traditional risk factors for HCC among HIV-infected patients. We hypothesized that higher HIV RNA levels, longer duration of HIV viremia, and lower CD4+ cell percentage, which is not affected by liver disease progression (13), were statistically significant determinants of HCC. We also examined the risk of HCC with traditional risk factors in the general population, including older age, black race, overweight or obesity, diabetes mellitus, alcohol dependence or abuse, tobacco use, and HBV and HCV coinfection (16). To account for the role of cirrhosis in the development of HCC and the possibility that risk factors for HCC might vary by presence of cirrhosis, we stratified our analysis by baseline cirrhosis status.

Methods

Study Design and Data Source

We conducted a retrospective cohort study among HIV-infected individuals in the Veterans Aging Cohort Study (VACS) between October 1, 1999, and September 30, 2015 (17). The VACS consists of electronic medical record data from HIV-infected patients receiving care at Veterans Health Administration (VA) facilities across the United States. Data include demographics, hospital and outpatient diagnoses (recorded using International

Classification of Diseases, Ninth Revision [ICD-9] codes), procedures, laboratory results, and dispensed medications. Death date was determined from the VA Vital Status File. The study was approved by the institutional review boards of the University of Pennsylvania, Corporal Michael J. Crescenz VA Medical Center in Philadelphia, VA Connecticut Healthcare System, and Yale University.

Study Patients

HIV-infected patients were included if they had HIV RNA, CD4+ cell count and percentage, and CD8+ cell count and percentage simultaneously assessed (which occurs routinely as part of HIV care in the VA system) between October 1, 1999, and September 30, 2015, and had at least 180 days of observation after determination of these laboratory results. We defined the start of follow-up as 180 days after the date that HIV RNA, CD4+, and CD8+ results were assessed. The 180 days prior to start of follow-up represented the baseline period, during which baseline comorbidities and laboratory results were collected. Patients were excluded if they had HCC diagnosed prior to start of follow-up. Follow-up continued until HCC, death, or last VA visit before September 30, 2015.

Main Study Outcomes

The primary outcome was incident HCC diagnosis. HCC diagnoses were determined from the VA national cancer registry by topography code C22.0 (liver) and morphology codes 8170–8180 (HCC) from the International Classification of Diseases for Oncology, Third Edition (ICD-O-3) (18), consistent with Surveillance, Epidemiology, and End Results coding algorithms (19). The VA cancer registry records cancers diagnosed and/or treated within the VA (7). ICD-O-3 codes validly identify cancer diagnoses, including HCC (20). To account for lags in reporting diagnoses in the cancer registry and minimize the likelihood of missing HCC events, we supplemented HCC case finding with ICD-9 diagnoses for HCC (155.0, 155.1, and 155.2) recorded in the VA electronic medical record. Prior research has shown that use of both VA cancer registry records and ICD-9 diagnoses have 90% sensitivity for incident cancer diagnosis when compared to chart review; however, positive predictive value varied (96% for VA cancer registry; 63% for ICD-9 diagnosis) (20,21). Consequently, HCC diagnoses from the registry and claims were confirmed by medical record review by trained adjudicators.

For all confirmed HCC diagnoses, we determined the presence of cirrhosis by review of medical records within 1 year prior to HCC diagnosis. Details on cirrhosis adjudication appear in the [Supplementary Methods](#) (available online).

Data Collection

Baseline data included age, sex, race and/or ethnicity, body mass index, diabetes [defined by random glucose ≥ 200 mg/dL, hemoglobin A1c $\geq 6.5\%$, or antidiabetic drug use (22)], alcohol dependence or abuse, injection and/or noninjection drug use, tobacco use (ever), HBV coinfection (ever positive HBV surface antigen), HCV coinfection (ever detectable HCV RNA or genotype), cirrhosis, HIV RNA, absolute CD4+ and CD8+ counts and percentages, and antiretroviral therapy (ART) use. Cirrhosis was defined by a hospital discharge diagnosis or outpatient diagnosis for cirrhosis or hepatic decompensation ([Supplementary Table 1](#) available online). Prior studies validated this determination within the VA system, with no less than 90% of these diagnoses confirmed by medical records (23,24). Alanine aminotransferase (ALT), aspartate aminotransferase (AST), and platelet count were collected from dates closest, but within 360 days prior, to start of follow-up. FIB-4, a noninvasive measure of hepatic fibrosis, was calculated by (age [years] \times AST [U/L]) / (platelet count [10^9 /L]) \times (ALT [U/L])^{1/2} (25).

Time-varying variables were assessed on a monthly basis and included HIV RNA, CD4+ cell count or percentage, CD8+ cell count or percentage, and diabetes. When multiple results of the same test were measured within a month, to be most conservative, we used the highest HIV RNA and CD8+ results and lowest CD4+ result. When a test was not updated during a month, we carried forward the value from the previous month until the next available result. HIV RNA, CD4+, and CD8+ results were updated at the beginning of each 30-day interval from baseline but were lagged by 180 days [approximate mean doubling time of HCC tumors <5 centimeters in length (26)] to reduce the possibility that the presence of HCC influenced these variables (ie, reverse causality).

Statistical Analysis

We determined unadjusted incidence rates (IR) of HCC (events per 100 000 person-years), overall and by HBV and HCV status. We used multivariable Cox regression to determine adjusted hazard ratios (HR; 95% confidence interval [CI]) of HCC for risk factors of interest. HIV-related factors included higher time-updated HIV RNA level, longer duration of detectable HIV (≥ 500 copies/mL), and lower time-updated CD4+ percentage. Traditional HCC risk factors examined included older age, black race, overweight or obesity, diabetes, alcohol dependence or abuse, ever use of tobacco, HBV coinfection, and HCV coinfection (16). Given the importance of cirrhosis on development of HCC, analyses were stratified by baseline cirrhosis status.

To evaluate the effects of HIV viremia on HCC risk, we created four separate models that examined HIV as a time-updated: continuous variable by 1.0 \log_{10} increments (model #1); categorical variable defined by detectable HIV (≥ 500 copies/mL; model #2); categorical variable classified by increasing categories of HIV RNA (<500 copies/mL; 500–9999 copies/mL; $\geq 10\,000$ copies/mL; model #3); and categorical variable classified by increasing consecutive months of detectable HIV (compared to those with undetectable HIV; model #4). For model #4, detectable HIV was evaluated as a monthly time-updated variable.

Once a patient was classified with HIV viremia, consecutive months were counted until a viral load less than 500 copies/mL was identified. If detectable HIV recurred, the count of consecutive months of viremia was restarted at one month. We determined hazard ratios of HCC associated with increasing consecutive months of detectable HIV (1–11 months; ≥ 12 months) compared to persons whose HIV RNA were suppressed throughout follow-up, adjusting for all other risk factors.

We performed five sensitivity analyses to assess the robustness of our results. First, we repeated the analysis accounting for competing risk of death (27). Second, we repeated analyses, lagging HIV RNA and CD4+ cell percentages by 360 and 540 days. Third, we evaluated risk factors for HCC separately among persons with HBV coinfection, HCV coinfection, and without viral hepatitis. Fourth, to explore the potential impact of hepatic steatosis on HCC risk, we classified patients with possible fatty liver disease prior to start of follow-up based on the presence of both obesity (body mass index ≥ 30 kg/m²) and diabetes mellitus, because these act synergistically to promote steatosis (28), and examined associations with HCC. Fifth, because duration of HIV-related immunosuppression might affect HCC risk, we determined whether longer consecutive months with CD4+ percentage less than 14% increased risk of HCC. Finally, in an exploratory analysis, we examined the risk of HCC with lower time-updated CD4+ to CD8+ ratio, which indicates dysfunctional immune activation (29), by cirrhosis status.

Proportionality of hazards was assessed by log-log plots and Schoenfeld residuals (30). To address the potential bias of missing data among covariates, we implemented multiple imputation using chained equations by means of 10 imputations using all variables in [Table 1](#) (31). Results across the 10 datasets were combined to arrive at confidence intervals that accounted for within- and across-dataset variances. Data were analyzed using SAS 9.4 (SAS Institute Inc., Cary, NC). All P values were two-sided. Statistical significance was declared if the 95% confidence interval did not cross 1.00.

Results

Patient Characteristics

We identified 37 946 HIV-infected individuals in the VACS who had HIV RNA, CD4+, and CD8+ results measured simultaneously between October 1, 1999, and September 30, 2015. After exclusions, 35 659 remained in the final sample.

Patients in the cohort had a median age of 46 years at baseline and were predominantly male, black, and overweight or obese ([Table 1](#)). Tobacco use, alcohol dependence or abuse, and injection and/or noninjection drug use were common. At the start of follow-up, 56.7% had HIV RNA level of 500 or more copies/mL, 23.9% had CD4+ percentage less than 14%, and 88.5% had a CD4+ to CD8+ ratio less than 1.0. A total of 31.9% had HCV coinfection, and 5.6% were HBV-coinfected. Only 11.9% of HCV-coinfected persons received anti-HCV therapy at baseline or during follow-up.

Overall, 68.7% received ART during the baseline period. ART regimens prescribed reflected the antiretrovirals used at the time of study entry ([Table 1](#)). Among 1981 HBV-coinfected individuals, 1211 (61.1%) were on ART at baseline. Of these, 689 (56.9%) received HBV-active ART with lamivudine or emtricitabine alone, 374 (30.9%) with tenofovir plus emtricitabine or lamivudine, and 27 (2.2%) with tenofovir alone; 121 (10.0%) were

Table 1. Baseline characteristics of study patients, stratified by cirrhosis diagnosis

Characteristic*	Overall No. (%) (n = 35 659)	Baseline no cirrhosis† No. (%) (n = 34 886)	Baseline cirrhosis‡ No. (%) (n = 733)
Median age (IQR), y	46 (39–53)	46 (39–53)	49 (44–55)
Male sex	34 792 (97.6)	34 032 (97.6)	760 (98.3)
Race/ethnicity			
Black	17 069 (47.9)	16 750 (48.0)	319 (41.3)
Caucasian	13 859 (38.9)	13 522 (38.8)	337 (43.6)
Hispanic	2720 (7.6)	2632 (7.5)	88 (11.4)
Other/Unknown	2011 (5.6)	1982 (5.7)	29 (3.8)
Body mass index			
Underweight, <18.50 kg/m ²	837 (2.3)	819 (2.3)	18 (2.3)
Normal, 18.50–24.99 kg/m ²	14 083 (39.5)	13 768 (39.5)	315 (40.8)
Overweight, 25.00–29.99 kg/m ²	11 415 (32.0)	11 184 (32.1)	231 (29.9)
Obesity, 30.00–34.99 kg/m ²	3882 (10.9)	3789 (10.9)	93 (12.0)
Morbid obesity, ≥35.00 kg/m ²	1323 (3.7)	1299 (3.7)	24 (3.1)
Missing weight and/or height	4119 (11.6)	4027 (11.5)	92 (11.9)
Diabetes mellitus	3308 (9.3)	3150 (9.0)	158 (20.4)
History of alcohol dependence/abuse	10 538 (29.6)	10 064 (28.8)	474 (61.3)
History of injection/noninjection drug use	16 235 (45.5)	15 781 (45.2)	454 (58.7)
Tobacco use			
Never	9533 (26.7)	9393 (26.9)	140 (18.1)
Ever‡	24 707 (69.3)	24 158 (69.2)	549 (71.0)
Unknown	1419 (4.0)	1335 (3.8)	84 (10.9)
Hepatitis C virus coinfection§			
Detectable HCV RNA or genotype	11 392 (31.9)	10 940 (31.4)	452 (58.5)
Ever treated with HCV antiviral	1354 (11.9)	1304 (11.9)	50 (11.1)
HCV antibody+/HCV RNA-	1055 (3.0)	1020 (2.9)	35 (4.5)
HCV antibody-	21 472 (60.2)	21 247 (60.9)	225 (29.1)
Never tested	1740 (4.9)	1679 (4.8)	61 (7.9)
Hepatitis B virus coinfection			
HBsAg+	1981 (5.6)	1873 (5.4)	108 (14.0)
Ever treated with HBV-active antiretroviral	1840 (92.9)	1748 (93.3)	92 (85.2)
HBsAg-	31 712 (88.9)	31 096 (89.1)	616 (79.7)
Never tested	1966 (5.5)	1917 (5.5)	49 (6.3)
HIV RNA			
Median (IQR), log ₁₀ cells/mm ³	3.2 (1.7–4.6)	3.2 (1.7–4.6)	3.0 (1.7–4.6)
≥500 copies/mL	20 216 (56.7)	19 791 (56.7)	425 (55.0)
CD4+ cell percentage			
Median (IQR)	22 (14–31)	22 (14–31)	22 (14–31)
≥28%	11 776 (33.0)	11 530 (33.1)	246 (31.8)
14–27.99%	14 798 (41.5)	14 456 (41.4)	342 (44.2)
<14%	8513 (23.9)	8337 (23.9)	176 (22.8)
Unknown	572 (1.6)	563 (1.6)	9 (1.2)
CD4+/CD8+ ratio			
Median (IQR)	0.40 (0.21–0.69)	0.40 (0.21–0.69)	0.42 (0.21–0.70)
<1.0	31 553 (88.5)	30 879 (88.5)	674 (87.2)
Median alanine aminotransferase (IQR), U/L	31 (21–47)	30 (21–47)	41 (27–71)
Not assessed at baseline	2362 (6.6)	2338 (6.7)	24 (3.1)
Median aspartate aminotransferase (IQR), U/L	30 (23–44)	29 (22–43)	55 (32–94)
Not assessed at baseline	1987 (5.6)	1964 (5.6)	23 (3.0)
Platelet count, x 10 ⁶ /L			
≥150 000	30 570 (85.7)	30 194 (86.6)	376 (48.6)
<150 000	4812 (13.5)	4420 (12.7)	392 (50.7)
Not assessed at baseline	277 (0.8)	272 (0.8)	5 (0.6)
Median baseline FIB-4 (IQR)	1.18 (0.81–1.78)	1.17 (0.81–1.74)	3.00 (1.58–5.82)
Unable to be calculated at baseline	3325 (9.3)	3290 (9.4)	35 (4.5)
On antiretroviral therapy	24 506 (68.7)	23 980 (68.7)	526 (68.0)
Most common baseline antiretroviral regimens¶			
Efavirenz/tenofovir/emtricitabine	3639 (14.8)	3565 (14.9)	74 (14.1)
Efavirenz/zidovudine/lamivudine	1764 (7.2)	1738 (7.2)	26 (4.9)
Nelfinavir/zidovudine/lamivudine	1177 (4.8)	1156 (4.8)	21 (4.0)
Indinavir/zidovudine/lamivudine	1033 (4.2)	1015 (4.2)	18 (3.4)

(continued)

Table 1. (continued)

Characteristic*	Overall No. (%) (n = 35 659)	Baseline no cirrhosis† No. (%) (n = 34 886)	Baseline cirrhosis‡ No. (%) (n = 733)
Atazanavir/tenofovir/emtricitabine	965 (3.9)	950 (4.0)	15 (2.9)
Nelfinavir/stavudine/lamivudine	888 (3.6)	863 (3.6)	25 (4.8)
Indinavir/stavudine/lamivudine	725 (3.0)	708 (3.0)	17 (3.2)
Nevirapine/zidovudine/lamivudine	663 (2.7)	652 (2.7)	11 (2.1)
Efavirenz/stavudine/lamivudine	611 (2.5)	590 (2.5)	21 (4.0)

HBsAg = hepatitis B surface antigen; HBV = hepatitis B virus; HCV = hepatitis C virus; HIV = human immunodeficiency virus; IQR = interquartile range; RNA = ribonucleic acid.

*Results reported as n (%) unless otherwise specified.

†Cirrhosis defined as any diagnosis of compensated or decompensated cirrhosis prior to the start of follow-up.

‡Ever tobacco use includes current and prior tobacco use.

§Hepatitis C virus coinfection defined as positive quantitative HCV RNA (absolute value determinable or not), positive qualitative HCV RNA, or quantifiable HCV genotype at baseline or during follow-up.

||Hepatitis B virus coinfection defined as positive HBV surface antigen at baseline or during follow-up.

¶Percentages represent proportion of antiretroviral use among patients prescribed therapy.

on ART without an HBV-active antiretroviral. By the end of follow-up, 1840 (92.9%) received HBV-active ART.

A total of 773 (2.2%) patients had a baseline diagnosis of cirrhosis. These patients were older and more commonly had diabetes, alcohol dependence or abuse, and HCV or HBV coinfection compared to those without cirrhosis (Table 1).

Incidence Rates of HCC

Overall, 302 (0.8%) medical record-confirmed HCC diagnoses were identified over 281 441 person-years (IR = 107.3/100 000 person-years) with median duration of follow-up of 7.4 (interquartile range = 3.3–12.8) years. Rates were particularly high among the 1981 HBV-coinfected (IR = 350.4/100 000 person-years) and 11 392 HCV-coinfected (IR = 224.6/100 000 person-years) individuals.

Patients diagnosed with HCC had a high prevalence of HCV coinfection (82.8%), alcohol dependence or abuse (63.2%), overweight or obesity (47.7%), and HBV coinfection (18.9%; Table 2). Notably, 99 (32.8%) did not have cirrhosis at HCC diagnosis (Table 2).

Determinants of HCC, by Baseline Cirrhosis Status

Among individuals without baseline cirrhosis, higher HIV RNA in model #1 (HR = 1.25, 95% CI = 1.12 to 1.40, per 1.0 log₁₀ copies/mL increase), detectable HIV in model #2 (HR 1.46, 95% CI = 1.07 to 1.99), HIV RNA of 10 000 or more copies/mL in model #3 (HR = 1.63, 95% CI = 1.11 to 2.40), and 12 or more months of detectable HIV viremia in model #4 (HR = 1.47, 95% CI = 1.02 to 2.11) increased the risk of HCC (Table 3). Absolute CD4+ counts less than 200 cells/mm³ increased HCC risk (Supplementary Table 2, available online); however, lower CD4+ percentage was not associated with an increased risk of HCC (Table 3). Additionally, in all models, older age, diabetes, HBV coinfection, HCV coinfection, history of alcohol dependence or abuse, and ever use of tobacco increased the risk of HCC (Table 3). Among individuals with baseline cirrhosis, only HCV coinfection was associated with an increased risk of HCC (Supplementary Table 3, available online).

Results were similar in analyses accounting for the competing risk of death (data not shown) and using 360-day and 540-day lags for HIV RNA and CD4+ cell percentage (Supplementary

Tables 4–5, available online). Similar findings were observed when analyses were restricted to HBV-coinfected (Supplementary Table 6, available online) and HCV-coinfected (Supplementary Table 7, available online) individuals, although results for some risk factors did not achieve statistical significance given the smaller sample sizes in these groups. There were too few HCC events (n = 17) among HIV-infected patients without viral hepatitis to permit analysis. When possible fatty liver disease was evaluated, patients who had both baseline obesity and diabetes had a higher risk of HCC than those who did not (HR = 2.32, 95% CI = 1.19 to 4.52; Supplementary Table 8, available online). Longer consecutive months with CD4+ percentage less than 14% did not increase HCC risk (≥12 months: HR = 1.02, 95% CI = 0.69 to 1.51; 1–11 months: HR = 1.15, 95% CI = 0.65 to 2.05) compared to those with no less than 14% throughout follow-up.

In exploratory analyses, after adjustment for HIV RNA and traditional risk factors, the risk of HCC was not increased with lower time-updated CD4+ to CD8+ ratio among either individuals with baseline cirrhosis (HR = 1.000, 95% CI = 0.999 to 1.001, per 0.1 unit decrease) or without cirrhosis (HR = 0.991, 95% CI = 0.981 to 1.002, per 0.1 unit decrease).

Discussion

To our knowledge, this is the largest study to evaluate HIV-related and traditional risk factors for HCC among HIV-infected patients and the first to examine such determinants by baseline cirrhosis status. We stratified our analyses by baseline cirrhosis status to account for the possibility that risk factors for HCC might vary by the presence of cirrhosis. Among patients without baseline cirrhosis, time-updated detectable HIV (≥500 copies/mL), higher HIV RNA (particularly ≥10 000 copies/mL), and no less than 12 months of detectable HIV increased risk of HCC, independent of traditional risk factors. Older age, HBV, HCV, diabetes, alcohol dependence or abuse, and tobacco use, which are traditional risk factors for HCC, also increased HCC risk in this group. The risk of HCC was particularly high in those with both obesity and diabetes. Among patients with baseline cirrhosis, only HCV coinfection remained associated with HCC. Notably, lower CD4+ cell percentage was not associated with increased risk of HCC regardless of cirrhosis status.

Our study is the first to find that higher level and longer duration of HIV viremia contribute to HCC risk. HIV viremia could

Table 2. Characteristics of patients with incident hepatocellular carcinoma, stratified by cirrhosis as determined by review of medical records within one year prior to diagnosis

Characteristic*	Overall incident HCC (n = 302)	No evidence of cirrhosis (n = 99)	Evidence of cirrhosis (n = 203)
Median age at diagnosis (IQR), y	56.4 (51.3–61.1)	57.5 (51.3–61.8)	56.0 (51.3–60.9)
Median time to diagnosis (IQR), y	7.2 (3.9–10.6)	6.3 (3.5–10.3)	7.4 (4.2–11.1)
Male sex	299 (99.0)	99 (100.0)	200 (98.5)
Race/ethnicity			
Black	159 (52.6)	62 (62.6)	97 (47.8)
Caucasian	97 (32.1)	24 (24.2)	73 (36.0)
Hispanic	37 (12.3)	10 (10.1)	27 (13.3)
Other/Unknown	9 (3.0)	3 (3.0)	6 (3.0)
Body mass index			
Underweight, <18.50 kg/m ²	25 (8.3)	16 (16.2)	9 (4.4)
Normal, 18.50–24.99 kg/m ²	128 (42.4)	47 (47.5)	81 (39.9)
Overweight, 25.00–29.99 kg/m ²	102 (33.8)	25 (25.3)	77 (37.9)
Obesity, 30.00–34.99 kg/m ²	34 (11.3)	9 (9.1)	25 (12.3)
Morbid obesity, ≥35.00 kg/m ²	8 (2.6)	2 (2.0)	6 (3.0)
Missing weight and/or height	5 (1.7)	0 (0.0)	5 (2.5)
Diabetes mellitus	94 (31.1)	24 (24.2)	70 (34.5)
History of alcohol dependence/abuse	191 (63.2)	61 (61.6)	130 (64.0)
History of injection/noninjection drug use	246 (81.5)	81 (81.8)	165 (81.3)
Tobacco use			
Never	40 (13.2)	10 (10.1)	30 (14.8)
Ever†	262 (86.8)	89 (89.9)	173 (85.2)
Hepatitis C virus coinfection‡			
Detectable HCV RNA or genotype	250 (82.8)	79 (79.8)	171 (84.2)
HCV antibody+/HCV RNA-	5 (1.7)	1 (1.0)	4 (2.0)
HCV antibody-	39 (12.9)	15 (15.2)	24 (11.8)
Never tested	8 (2.6)	4 (4.0)	4 (2.0)
Hepatitis B virus coinfection§			
HBsAg+	57 (18.9)	17 (17.2)	40 (19.7)
HBsAg-	236 (78.1)	79 (79.8)	157 (77.3)
Never tested	9 (3.0)	3 (3.0)	6 (3.0)
Median HIV RNA (IQR), log ₁₀ copies/mL	1.7 (1.7–2.6)	1.7 (1.7–2.6)	1.7 (1.7–2.7)
CD4 percentage			
Median (IQR), %	26 (17–35)	26 (14–35)	26 (18–35)
≥28%	135 (44.7)	42 (42.4)	93 (45.8)
14–27.99%	120 (39.7)	35 (35.4)	85 (41.9)
<14%	47 (15.6)	22 (22.2)	25 (12.3)
CD4: CD8 ratio			
Median (IQR)	0.57 (0.31–0.93)	0.52 (0.26–0.98)	0.59 (0.33–0.91)
<1.0	239 (79.1)	77 (77.8)	162 (79.8)
Median alanine aminotransferase (IQR), U/L	54 (36–79)	52 (35–74)	57 (36–79)
Not assessed within 360 days prior to HCC diagnosis	2 (0.7)	0 (0.0)	2 (1.0)
Median aspartate aminotransferase (IQR), U/L	68 (44–101)	61 (38–95)	73 (47–104)
Not assessed within 360 days prior to HCC diagnosis	1 (0.3)	0 (0.0)	1 (0.5)
Platelet count <150 000 × 10 ⁶ /L	172 (57.0)	42 (42.4)	130 (64.0)
FIB-4			
<1.45	27 (8.9)	18 (18.2)	9 (4.4)
1.45–3.25	99 (32.8)	39 (39.4)	60 (29.6)
>3.25	173 (57.3)	42 (42.4)	131 (64.5)
Insufficient data to calculate FIB-4	3 (1.0)	0 (0.0)	3 (1.5)
On antiretroviral therapy	229 (75.8)	77 (77.8)	152 (74.9)
Ever dideoxynucleoside analogue use¶	207 (68.5)	74 (74.7)	133 (65.5)

HBsAg = hepatitis B surface antigen; HCC = hepatocellular carcinoma; HCV = hepatitis C virus; IQR = interquartile range; NRTI = nucleoside reverse transcriptase inhibitors; RNA = ribonucleic acid.

*Results reported as n (%) unless otherwise specified.

†Ever tobacco use includes current and prior tobacco use.

‡Hepatitis C virus coinfection defined as positive quantitative HCV RNA (absolute value determinable or not), positive qualitative HCV RNA, or quantifiable HCV genotype at baseline or during follow-up.

§Hepatitis B virus coinfection defined as positive HBV surface antigen at baseline or during follow-up.

||FIB-4 was calculated using current age and most recent alanine aminotransferase, aspartate aminotransferase, and platelet count within 360 days prior to HCC diagnosis.

¶Included didanosine, stavudine, zalcitabine, and zidovudine.

Table 3. Factors associated with incident hepatocellular carcinoma among HIV-infected patients in the Veterans Aging Cohort Study (October 1, 1999–September 30, 2015) without a baseline diagnosis of cirrhosis (n = 34 886; 270 hepatocellular carcinoma events)

Characteristic	Unadjusted HR (95% CI)	Model #1* Adj. HR (95% CI)	Model #2† Adj. HR (95% CI)	Model #3‡ Adj. HR (95% CI)	Model #4§ Adj. HR (95% CI)
Age, per 10 y	1.33 (1.17 to 1.51)	1.48 (1.26 to 1.73)	1.44 (1.23 to 1.69)	1.45 (1.23 to 1.70)	1.44 (1.23 to 1.69)
Male sex	2.30 (0.74 to 7.19)	1.38 (0.44 to 4.31)	1.37 (0.44 to 4.30)	1.37 (0.44 to 4.29)	1.37 (0.44 to 4.30)
Race	Reference	Reference	Reference	Reference	Reference
White	1.50 (1.15 to 1.97)	0.97 (0.73 to 1.28)	0.97 (0.74 to 1.29)	0.97 (0.74 to 1.29)	0.97 (0.74 to 1.29)
Black	1.77 (1.17 to 2.68)	1.24 (0.82 to 1.89)	1.23 (0.81 to 1.87)	1.23 (0.81 to 1.88)	1.23 (0.81 to 1.87)
Hispanic	1.40 (0.68 to 2.89)	1.72 (0.83 to 3.58)	1.72 (0.83 to 3.56)	1.71 (0.83 to 3.55)	1.72 (0.83 to 3.56)
Other					
Baseline body mass index					
Underweight, <18.50 kg/m ²	0.63 (0.18 to 2.15)	0.59 (0.17 to 2.08)	0.59 (0.17 to 2.09)	0.59 (0.17 to 2.09)	0.59 (0.17 to 2.09)
Normal, 18.50–24.99 kg/m ²	Reference	Reference	Reference	Reference	Reference
Overweight, 25.00–29.9 kg/m ²	0.88 (0.67 to 1.17)	0.96 (0.72 to 1.27)	0.96 (0.72 to 1.27)	0.96 (0.72 to 1.27)	0.96 (0.72 to 1.27)
Obesity, 30.00–34.9 kg/m ²	0.92 (0.61 to 1.40)	1.01 (0.65 to 1.56)	1.01 (0.65 to 1.56)	1.01 (0.65 to 1.56)	1.01 (0.65 to 1.56)
Morbid obesity, ≥35.00 kg/m ²	0.84 (0.39 to 1.80)	0.98 (0.45 to 2.14)	0.98 (0.45 to 2.13)	0.98 (0.45 to 2.13)	0.98 (0.45 to 2.13)
Time-updated diabetes mellitus	1.70 (1.32 to 2.20)	1.46 (1.12 to 1.91)	1.45 (1.11 to 1.90)	1.45 (1.11 to 1.90)	1.45 (1.11 to 1.90)
Hepatitis B virus coinfection	Reference	Reference	Reference	Reference	Reference
HBsAg-	3.65 (2.68 to 4.97)	3.92 (2.87 to 5.35)	3.91 (2.86 to 5.34)	3.91 (2.86 to 5.35)	3.91 (2.86 to 5.34)
HBsAg+	1.01 (0.48 to 2.15)	1.12 (0.52 to 2.42)	1.13 (0.52 to 2.44)	1.13 (0.52 to 2.44)	1.13 (0.52 to 2.44)
Never tested					
Hepatitis C virus coinfection	Reference	Reference	Reference	Reference	Reference
HCV antibody-	9.25 (6.53 to 13.11)	7.65 (5.35 to 10.94)	7.68 (5.36 to 10.98)	7.68 (5.37 to 11.00)	7.68 (5.36 to 10.98)
Detectable HCV RNA or genotype	4.83 (1.90 to 12.28)	3.73 (1.46 to 9.52)	3.81 (1.49 to 9.72)	3.80 (1.49 to 9.71)	3.81 (1.49 to 9.72)
HCV antibody+/HCV RNA-	4.85 (2.16 to 10.88)	4.64 (2.04 to 10.55)	4.68 (2.06 to 10.65)	4.68 (2.06 to 10.64)	4.68 (2.06 to 10.65)
Never tested					
Time-updated CD4+ cell percentage	Reference	Reference	Reference	Reference	Reference
≥28%	1.05 (0.81 to 1.37)	0.86 (0.66 to 1.12)	0.89 (0.68 to 1.16)	0.88 (0.68 to 1.16)	0.89 (0.68 to 1.16)
14–27.99%	1.52 (1.08 to 2.14)	0.97 (0.66 to 1.43)	1.12 (0.78 to 1.62)	1.09 (0.75 to 1.59)	1.12 (0.78 to 1.62)
<14%	2.38 (1.85 to 3.05)	1.45 (1.11 to 1.89)	1.46 (1.12 to 1.90)	1.46 (1.12 to 1.90)	1.46 (1.12 to 1.90)
History of alcohol abuse	Reference	Reference	Reference	Reference	Reference
Tobacco use	2.58 (1.81 to 3.67)	1.65 (1.14 to 2.38)	1.66 (1.15 to 2.39)	1.66 (1.15 to 2.40)	1.66 (1.15 to 2.39)
Never	1.26 (1.14 to 1.39)	1.25 (1.12 to 1.40)	—	—	—
Ever	1.58 (1.19 to 2.10)	—	1.46 (1.07 to 1.99)	—	—
Time-updated HIV RNA, per 1.0 log ₁₀ copies/mL	Reference	Reference	Reference	Reference	Reference
Current HIV RNA ≥500 copies/mL	1.41 (0.94 to 2.11)	—	—	1.31 (0.87 to 1.97)	—
Current HIV RNA <500 copies/mL	1.73 (1.22 to 2.47)	—	—	1.63 (1.11 to 2.40)	—

(continued)

Table 3. (continued)

Characteristic	Unadjusted HR (95% CI)	Model #1* Adj. HR (95% CI)	Model #2† Adj. HR (95% CI)	Model #3‡ Adj. HR (95% CI)	Model #4§ Adj. HR (95% CI)
Consecutive months of HIV RNA \geq 500 copies/mL	Reference	—	—	—	Reference
HIV RNA always < 500 copies/mL	1.63 (1.07 to 2.49)	—	—	—	1.46 (0.94 to 2.26)
1–11 months of HIV RNA \geq 500 copies/mL	1.55 (1.10 to 2.18)	—	—	—	1.47 (1.02 to 2.11)
\geq 12 months of HIV RNA \geq 500 copies/mL					

CI = confidence interval; HBsAg = hepatitis B surface antigen; HBV = hepatitis B virus; HCV = hepatitis C virus; HR = hazard ratio; RNA = ribonucleic acid.

*Model #1 includes continuous values of HIV RNA as log₁₀ copies/mL.

†Model #2 includes current HIV RNA \geq 500 copies/mL.

‡Model #3 includes current categories of HIV RNA \geq 500 copies/mL.

§Model #4 includes consecutive months of HIV RNA \geq 500 copies/mL.

||Ever tobacco use includes current and prior tobacco use.

¶Two-sided $P_{trend} = .01$, based on multivariable Cox regression analysis.

contribute to HCC by accelerating hepatic fibrosis progression to cirrhosis or by directly promoting hepatocarcinogenesis via immune dysregulation, oxidative stress, hepatocyte apoptosis, and/or depletion of CD4+ cells in the gastrointestinal tract with resultant microbial translocation (32–34). We have previously shown that suppression of HIV viremia can delay onset of cirrhosis (21). Our findings suggest that achieving and maintaining HIV suppression could mitigate the risk of HCC.

Three prior studies found no association between HIV RNA and risk of HCC, but none stratified analyses by baseline cirrhosis status. One study of 31 576 HIV-infected patients in the VA HIV Clinical Case Registry from 1985 to 2010 found that longer percentage of time with undetectable HIV RNA (<500 copies/mL) did not decrease HCC risk (5). A follow-up study among 8563 HIV and HCV-coinfected patients in this registry similarly found no association between duration of undetectable HIV RNA and HCC (6). However, both analyses included cirrhosis as a covariate in multivariable models. Because cirrhosis is in the causal pathway to HCC, controlling for cirrhosis could have adjusted away associations between HIV suppression and HCC. A third study among 42 441 HIV-infected patients in the VACS from 1999 to 2015 found that neither early (\leq 2 years) nor long-term ($>$ 2 years) HIV suppression decreased rates of HCC compared to 104 712 demographically similar uninfected persons (7). However, this analysis did not stratify results by baseline cirrhosis status.

Contrary to previous studies (6,8–12), we found that HIV-related immunosuppression, as measured by CD4+ cell percentage, was not associated with an increased risk of HCC. Notably, those prior studies evaluated the risk of HCC associated with lower absolute CD4+ count. Indeed, when we evaluated associations between absolute CD4+ count and HCC in our cohort, CD4+ counts less than 200 cells/mm³ increased HCC risk. However, absolute CD4+ count may decrease during cirrhosis as a result of portal hypertension-induced splenic sequestration, but CD4+ percentage remains unchanged during cirrhosis (13). Our results suggest that HIV-related immunosuppression is not an important contributor to HCC risk and that the findings of prior analyses likely reflected the effect of liver fibrosis progression on absolute CD4+ count.

Interestingly, 32.8% of patients with HCC in our study did not have evidence of cirrhosis based on review of medical records within one year prior to cancer diagnosis. HCC can develop in the absence of advanced hepatic fibrosis in chronic HBV infection or nonalcoholic fatty liver disease (35,36). Further research is needed to determine how frequently HCC occurs in the absence of cirrhosis in HIV and if this differs from uninfected persons.

The study has several potential limitations. First, we might have underestimated cirrhosis, because this condition is clinically silent. However, cirrhosis was identified using validated diagnoses, and the negative predictive value of this definition exceeded 99% (23,24). Second, we were unable to determine fatty liver disease, because this diagnosis requires liver imaging or biopsy to confirm. We classified possible fatty liver disease by baseline presence of both obesity and diabetes. Future studies should evaluate the effect of fatty liver disease on HCC in HIV. Third, we did not evaluate the risk of HCC with antiretroviral drugs, particularly those associated with hepatotoxicity (37), or viral hepatitis treatments. Additional research should evaluate the effects of these medications on incidence of HCC. Finally, our sample was predominantly comprised of male US veterans, but the study included 867 HIV-infected women.

In conclusion, among HIV-infected patients without cirrhosis, higher HIV RNA and longer duration of HIV viremia, in addition to HBV and HCV coinfection, were important determinants

of HCC, independent of traditional risk factors. HIV-related immunosuppression, determined by CD4+ cell percentage, was not associated with increased risk of HCC. This study provides the strongest evidence to date that HIV viremia contributes to the risk of HCC in this group.

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Affiliations of authors: Division of Infectious Diseases, Department of Medicine (JT, VLR), and Department of Biostatistics, Epidemiology, and Informatics, Center for Clinical Epidemiology and Biostatistics, Center for Pharmacoepidemiology Research and Training (JT, MJK, DMC, VLR), Perelman School of Medicine, University of Pennsylvania, Philadelphia, PA; Center for Population Health Sciences, Stanford University School of Medicine, Stanford, CA (LSP); VA Connecticut Healthcare System, West Haven, CT (RLM, KD, JPT, JKL, THT, ACJ); Yale University School of Medicine, New Haven, CT (RLM, KD, JPT, JKL, THT, ACJ); VA Greater Los Angeles Healthcare System and David Geffen School of Medicine at UCLA, Los Angeles, CA (MBG); Infectious Diseases Section, Michael E. DeBakey VA Medical Center and Department of Medicine, Baylor College of Medicine, Houston, TX (MCR); Washington DC VA Medical Center and George Washington University Medical Center, Washington, DC (CLG); James J. Peters VA Medical Center, Bronx, NY, and Icahn School of Medicine at Mount Sinai, New York, NY (NB, STB); Department of Biostatistics, Rutgers University School of Public Health, New Brunswick, NJ (JAR).

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